==========

Sequence Listing was accepted.

See attached Validation Report.

If you need help call the Patent Electronic Business Center at (866) 217-9197 (toll free).

Reviewer: Durreshwar Anjum

Timestamp: Wed Oct 17 12:50:35 EDT 2007

==========

Validated By CRFValidator v 1.0.3

Application No: 10595559 Version No: 1.0

Input Set:

Output Set:

Started: 2007-10-01 13:20:46.903
Finished: 2007-10-01 13:20:48.664
Elapsed: 0 hr(s) 0 min(s) 1 sec(s) 761 ms
Total Warnings: 16
Total Errors: 1
No. of SeqIDs Defined: 51
Actual SeqID Count: 51

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E	257	Invalid sequence data feature in <221> in SEQ ID (36)
W	213	Artificial or Unknown found in <213> in SEQ ID (37)
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SEQUENCE LISTING

<110> GUTHRIDGE, MARK
RAMSHAW, HAYLEY
STOMSKI, FRANK
LOPEZ, ANGEL

<120> A BINDING MOTIF AND METHODS OF REGULATING CELL FUNCTION

<130> 03391/0204241-US0

<140> 10595559
<141> 2007-10-01

<150> PCT/AU04/01480
<151> 2004-10-27

<150> AU 2003-905932
<151> 2003-10-27

<160> 51

<170> PatentIn Ver. 3.3

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Trp Glu Arg Ser Leu Ala Gly Ala Glu Glu Thr Ile Pro Leu Gln Thr
             20                  25                  30

Leu Arg Cys Tyr Asn Asp Tyr Thr Ser His Ile Thr Cys Arg Trp Ala
         35                  40                  45

Asp Thr Gln Asp Ala Gln Arg Leu Val Asn Val Thr Leu Ile Arg Arg
     50                  55                  60

Val Asn Glu Asp Leu Leu Glu Pro Val Ser Cys Asp Leu Ser Asp Asp
 65                  70                  75                  80

Met Pro Trp Ser Ala Cys Pro His Pro Arg Cys Val Pro Arg Arg Cys
                 85                  90                  95

Val Ile Pro Cys Gln Ser Phe Val Val Thr Asp Val Asp Tyr Phe Ser
            100                 105                 110

Phe Gln Pro Asp Arg Pro Leu Gly Thr Arg Leu Thr Val Thr Leu Thr
        115                 120                 125

Gln His Val Gln Pro Pro Glu Pro Arg Asp Leu Gln Ile Ser Thr Asp
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    130                 135                 140

Gln Asp His Phe Leu Leu Thr Trp Ser Val Ala Leu Gly Ser Pro Gln
145                 150                 155                 160

Ser His Trp Leu Ser Pro Gly Asp Leu Glu Phe Glu Val Val Tyr Lys
                165                 170                 175

Arg Leu Gln Asp Ser Trp Glu Asp Ala Ala Ile Leu Leu Ser Asn Thr
            180                 185                 190

Ser Gln Ala Thr Leu Gly Pro Glu His Leu Met Pro Ser Ser Thr Tyr
        195                 200                 205

Val Ala Arg Val Arg Thr Arg Leu Ala Pro Gly Ser Arg Leu Ser Gly
    210                 215                 220

Arg Pro Ser Lys Trp Ser Pro Glu Val Cys Trp Asp Ser Gln Pro Gly
225                 230                 235                 240

Asp Glu Ala Gln Pro Gln Asn Leu Glu Cys Phe Phe Asp Gly Ala Ala
                245                 250                 255

Val Leu Ser Cys Ser Trp Glu Val Arg Lys Glu Val Ala Ser Ser Val
            260                 265                 270

Ser Phe Gly Leu Phe Tyr Lys Pro Ser Pro Asp Ala Gly Glu Glu Glu
        275                 280                 285

Cys Ser Pro Val Leu Arg Glu Gly Leu Gly Ser Leu His Thr Arg His
    290                 295                 300

His Cys Gln Ile Pro Val Pro Asp Pro Ala Thr His Gly Gln Tyr Ile
305                 310                 315                 320

Val Ser Val Gln Pro Arg Arg Ala Glu Lys His Ile Lys Ser Ser Val
                325                 330                 335

Asn Ile Gln Met Ala Pro Pro Ser Leu Asn Val Thr Lys Asp Gly Asp
            340                 345                 350

Ser Tyr Ser Leu Arg Trp Glu Thr Met Lys Met Arg Tyr Glu His Ile
        355                 360                 365

Asp His Thr Phe Glu Ile Gln Tyr Arg Lys Asp Thr Ala Thr Trp Lys
    370                 375                 380

Asp Ser Lys Thr Glu Thr Leu Gln Asn Ala His Ser Met Ala Leu Pro
385                 390                 395                 400

Ala Leu Glu Pro Ser Thr Arg Tyr Trp Ala Arg Val Arg Val Arg Thr
                405                 410                 415

Ser Arg Thr Gly Tyr Asn Gly Ile Trp Ser Glu Trp Ser Glu Ala Arg
            420                 425                 430

Ser Trp Asp Thr Glu Ser Val Leu Pro Met Trp Val Leu Ala Leu Ile
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        435                 440                 445

Val Ile Phe Leu Thr Ile Ala Val Leu Leu Ala Leu Arg Phe Cys Gly
    450                 455                 460

Ile Tyr Gly Tyr Arg Leu Arg Arg Lys Trp Glu Glu Lys Ile Pro Asn
465                 470                 475                 480

Pro Ser Lys Ser His Leu Phe Gln Asn Gly Ser Ala Glu Leu Trp Pro
                485                 490                 495

Pro Gly Ser Met Ser Ala Phe Thr Ser Gly Ser Pro Pro His Gln Gly
            500                 505                 510

Pro Trp Gly Ser Arg Phe Pro Glu Leu Glu Gly Val Phe Pro Val Gly
        515                 520                 525

Phe Gly Asp Ser Glu Val Ser Pro Leu Thr Ile Glu Asp Pro Lys His
    530                 535                 540

Val Cys Asp Pro Pro Ser Gly Pro Asp Thr Thr Pro Ala Ala Ser Asp
545                 550                 555                 560

Leu Pro Thr Glu Gln Pro Pro Ser Pro Gln Pro Gly Pro Pro Ala Ala
                565                 570                 575

Ser His Thr Pro Glu Lys Gln Ala Ser Ser Phe Asp Phe Asn Gly Pro
            580                 585                 590

Tyr Leu Gly Pro Pro His Ser Arg Ser Leu Pro Asp Ile Leu Gly Gln
        595                 600                 605

Pro Glu Pro Pro Gln Glu Gly Gly Ser Gln Lys Ser Pro Pro Pro Gly
    610                 615                 620

Ser Leu Glu Tyr Leu Cys Leu Pro Ala Gly Gly Gln Val Gln Leu Val
625                 630                 635                 640

Pro Leu Ala Gln Ala Met Gly Pro Gly Gln Ala Val Glu Val Glu Arg
                645                 650                 655

Arg Pro Ser Gln Gly Ala Ala Gly Ser Pro Ser Leu Glu Ser Gly Gly
            660                 665                 670

Gly Pro Ala Pro Pro Ala Leu Gly Pro Arg Val Gly Gly Gln Asp Gln
        675                 680                 685

Lys Asp Ser Pro Val Ala Ile Pro Met Ser Ser Gly Asp Thr Glu Asp
    690                 695                 700

Pro Gly Val Ala Ser Gly Tyr Val Ser Ser Ala Asp Leu Val Phe Thr
705                 710                 715                 720

Pro Asn Ser Gly Ala Ser Ser Val Ser Leu Val Pro Ser Leu Gly Leu
                725                 730                 735

Pro Ser Asp Gln Thr Pro Ser Leu Cys Pro Gly Leu Ala Ser Gly Pro
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            740                 745                 750

Pro Gly Ala Pro Gly Pro Val Lys Ser Gly Phe Glu Gly Tyr Val Glu
        755                 760                 765

Leu Pro Pro Ile Glu Gly Arg Ser Pro Arg Ser Pro Arg Asn Asn Pro
    770                 775                 780

Val Pro Pro Glu Ala Lys Ser Pro Val Leu Asn Pro Gly Glu Arg Pro
785                 790                 795                 800

Ala Asp Val Ser Pro Thr Ser Pro Gln Pro Glu Gly Leu Leu Val Leu
                805                 810                 815

Gln Gln Val Gly Asp Tyr Cys Phe Leu Pro Gly Leu Gly Pro Gly Pro
            820                 825                 830

Leu Ser Leu Arg Ser Lys Pro Ser Ser Pro Gly Pro Gly Pro Glu Ile
        835                 840                 845

Lys Asn Leu Asp Gln Ala Phe Gln Val Lys Lys Pro Pro Gly Gln Ala
    850                 855                 860

Val Pro Gln Val Pro Val Ile Gln Leu Phe Lys Ala Leu Lys Gln Gln
865                 870                 875                 880

Asp Tyr Leu Ser Leu Pro Pro Trp Glu Val Asn Lys Pro Gly Glu Val
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Arg Tyr Phe Thr Gln Lys Glu Glu Thr Glu Ser Gly Ser Gly Pro
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<211> 22
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<213> Homo sapiens

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Asn Lys Lys Tyr Glu Leu Gln Asp Arg Asp Val Cys Glu Pro Arg Tyr
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Arg Ser Val Ser Glu Pro
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<211> 13
<212> PRT
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<400> 6
Asn Pro Thr Tyr Ser Val Met Arg Ser His Ser Tyr Pro
  1   5   10

<210> 7
<211> 24
<212> PRT
<213> Homo sapiens

<400> 7
Asn Ile Phe Tyr Leu Ile Arg Lys Ser Gly Ser Phe Pro Met Pro Glu
  1   5   10   15

Leu Lys Leu Ser Ile Ser Phe Pro
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<210> 8
<211> 19
<212> PRT
<213> Homo sapiens

<400> 8
Asn Glu Glu Tyr Leu Asp Leu Ser Gln Pro Leu Glu Gln Tyr Ser Pro
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Ser Tyr Pro

<210> 9
<211> 19

<212> PRT
<213> Homo sapiens

<400> 9
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1 5 10 15

Ser Phe Pro

<210> 10
<211> 16
<212> PRT
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<400> 10
Asn Ala Thr Tyr Lys Val Asp Val Ile Gln Arg Thr Arg Ser Lys Pro
1 5 10 15

<210> 11
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<400> 11
Asn Pro Glu Tyr His Ser Ala Ser Ser Gly Pro
1 5 10

<210> 12
<211> 10
<212> PRT
<213> Homo sapiens

<400> 12
Asn Pro Asp Tyr Trp Asn His Ser Leu Pro
1 5 10

<210> 13
<211> 23
<212> PRT
<213> Homo sapiens

<400> 13
Asn Pro Ser Tyr Ser Ser Asn Pro Phe Val Asn Tyr Asn Lys Thr Ser
1 5 10 15

Ile Cys Ser Lys Ser Asn Pro
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<210> 14
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<213> Homo sapiens

<400> 14
Asn Thr Leu Tyr Phe Asn Ser Gln Ser Ser Pro
  1   5   10

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<400> 15
Asn Pro Val Tyr Gln Lys Thr Thr Glu Asp Glu Val His Ile Cys His
  1   5   10   15

Asn Gln Asp Gly Tyr Ser Tyr Pro
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<210> 16
<211> 24
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Gly Arg His Ser Ala Ser Val Gly
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<210> 17
<211> 38
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Asn Pro Thr Tyr Lys Met Tyr Glu Gly Gly Glu Pro Asp Asp Val Gly
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Gly Leu Leu Asp Ala Asp Phe Ala Leu Asp Pro Asp Lys Pro Thr Asn
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Phe Thr Asn Pro Val Tyr
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Asn Pro Ile Tyr Lys Ser Ala Val Thr Thr Val Val
  1   5   10

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  1   5   10

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<400> 20
Asn Pro Leu Tyr Lys Glu Ala Thr Ser Thr Phe Thr
  1   5   10

<210> 21
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<400> 21
Asn Pro Leu Tyr Arg Lys Pro Ile Ser Thr His Thr
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<210> 22
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<400> 22
Asn Pro Leu Tyr Arg Gly Ser Thr Ser Thr Phe Lys
  1   5   10

<210> 23
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<400> 23
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<210> 24
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Val Gln Thr Tyr Val Leu Gln Gly Asp Pro Arg Ala Val Ser Thr Gln
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Pro

<210> 25
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Ser Glu Ser Thr Gln Pro
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Trp Lys Met Tyr Glu Val Tyr Asp Ala Lys Ser Lys Ser Val Ser Leu
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Pro

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<400> 28
Lys Ile Pro Tyr Phe His Ala Gly Gly Ser Lys Cys Ser Thr Trp Pro
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<210> 29
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<213> Homo sapiens

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Ser Pro Pro

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Asp Leu Glu Pro Glu Asn Met Glu Ser Val Pro Leu Asp Pro Ser Ala
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Ser Ser Ser Ser Leu Pro
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Lys Arg Pro Ser Phe Pro
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<400> 33
Ile Ser Gln Tyr Leu Gln Asn Ser Lys Arg Lys Ser Arg Pro
  1   5   10

<210> 34
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<400> 34
Gly Thr Ala Tyr Gly Leu Ser Arg Ser Gln Pro
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<210> 35
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<400> 35
Tyr Leu Pro Gln Glu Asp Trp Ala Pro Thr Ser Leu Thr Arg Pro
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20 25 30

Gly Glu Lys Leu His Ser Asp Ser Gly Ile Ser
35 40

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<210> 38

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<210> 39
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<223> Description of Artificial Sequence: Synthetic peptide

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